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Result
                                                                                                                                                                                                                                                                                                         Minimum DB
Maximum DB
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                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 0%
Maximum Match 10
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                                                                                                                                                                                                                                                                                                                                                                                                                       OM protein
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   and is derived by analysis of the total score distribution
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                                                                                                                                                                     Pred. No.
                                                                                                                          Score
                                                                                                                                                                                                                                                                                                         seq length: 0 seq length: 2000000000
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                                                                                                                                                            No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed,
   protein search, using sw model
                                                                                                                        %
Query
Match Length !
                                                                                                                                                                                                                                                         SPTREMBL_15:*
1: sp_archea:*
2: sp_bacteria
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9:
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11:
12:
13:
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  US-09-331-631A-37
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                                                                                                                                                                                                                                                                                                                                        374700 seqs, 117207915 residues
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Copyright (c) 1993 - 2000 Compugen Ltd
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   sp_mhc:*
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first 45 summaries
   2001, 16:22:07
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   Q9V110
Q9N9H2
P8219H2
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093593
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013259
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Q9VIY8
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P79377
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Q23947
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O9n9h1 ruditapes d
O92044 cyprinodon
O73914 notothenia
O93593 chaenocepha
O93609 trematomus
O13259 chionodraco
O13258 chionodraco
O13586 pagrus majo
O91910 cyprinus ca
                                                      Q23948 drosophila
Q9yi10 bungarus mu
Q9n9h2 ruditapes p
P82107 hirudo medi
                                                                                           Q16861 homo sapien
Q9viy8 drosophila
Q9rfp5 mycoplasma
P79377 sus scrofa
                                                                               P79380 sus scrofa
Q23947 drosophila
                                                                                                                          Description
                                                                                                             homo sapien
                                                                                                                                            RESULT
Q16861
                                                                                                Qy
                                                                                     DЬ
                                                                                                                                                                                                                   Q16861;
Q16861;
Q1-NOV-1996;
Q1-NOV-1996;
Q1-AUG-1998;
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ALIGNMENTS

(TrEMBLrel. 01, Created)
(TrEMBLrel. 01, Last sequence update)
(TrEMBLrel. 07, Last annotation update)

PRELIMINARY;

PRT;

46

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RESULT 2
Q9VIY8 PRELIN
AC Q9VIY8;
AC Q9VIY8;
DT 01-MAY-2000 (TrEME
DT 01-MAY-2000 (TREME
DT 01-MAY-2000 (TREME
DF 01-MAY-2000 (TREME
DE CG17567 PROTEIN.
GN CG17567.
OS Drosophila melanoc
OC Eukaryota; Metazoa
OC Pterygota; Neoptea
                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 4; Conserv
Drosophila |
Eukaryota; |
Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                         EMBL; U63332; AABOS810.1;
NON_TER 1 1 1
SEQUENCE 46 AA; 5061 MM
                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
Lemasson I., Devaux C., Mesnard J.M.;
Submitted (JUL-1996) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
                                                                                                                                                                                                                                        11
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                   melanogaster (Fruit fly).
Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
 Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                 (TrEMBLrel. 13, Created)
(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 13, Last annotation update)
                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                              5061 MW;
                                                                                                                                                                                                                                                                                                                       100.0%;
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RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,

RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottler P.,

RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Clandra I.,

RA Gerry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,

RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,

RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,

RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,

RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,

RA Hostin D., Houston K.A., Howland T.J., Wei M., H., Ibegwam C.,

RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,

RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,

RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,

RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,

RA Meinert K., Remington K.A., Nixon K., Nusskern D.R., Pacleb J.M.,

RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,

RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,

RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

RA Zheng X.H., Zhong F.N., Zhong W., Zhang G., Zhao Q., Zheng L.,

RA Zheng S.H., Zhong F.N., Zhong W., Zhan M., Shu S., Zhu X., Smith H.O.,

RA Zheng S.H., Zhong F.N., Zhong W., Zhong X., Zhu X., Smith H.O.,

RA Zheng S.H., Zhong G.Z., Sapsaj J.; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
SEQUENCE FROM N.A. STRAIN=PG18;
                                                                                   Mycoplasmataceae; Mycoplasma NCBI_TaxID=2115;
                                                                                                                                         Bacteria; Firmicutes; Bacillus/Clostridium group; Mollicutes;
                                                                                                                                                                          Mycoplasma
                                                                                                                                                                                                                                  01-MAY-2000
01-MAY-2000
                                                                                                                                                                                                                                                                                           Q9RFP5;
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                       Q9RFP5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=20196006; PubMed=10731132;
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SE; FBgn0040994; CG17567.
NCE 47 AA; 4909 MW; 1
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                                                                                                                                                                       fermentans.
                                                                                                                                                                                                                            (TrEMBLrel. 13, (TrEMBLrel. 13, (TrEMBLrel. 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                            Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                           Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
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79;
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Query Match
Best Local Similarity
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Best Local
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p79377;
01-MAY-1997 (TrEMBLrel. 03, C
01-MAY-1997 (TrEMBLrel. 03, L
01-OCT-2000 (TrEMBLrel. 15, L
METALLOTHIONEIN (MT) (FRAGMEN
                                                                   METAL
SEQUENCE
                                                                                          METAL
METAL
METAL
METAL
                                                                                                                                                                                                                                                                                                        Gene
                                                                                                                                                                                        PRINTS;
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HSSP; P02795; 1
                                                                                                                                                                                                                                                                                                                                                      MEDLINE-98241500;
                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence during insertion.";
J. Bacteriol 181:7597-7607(1999).
EMBL; AF179376; AAF15567.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Calcutt M.J., Lavrrar J.L., Wise K.S.; "IS1630 of Mycoplasma fermentans, a novel IS30-type insertion element that targets and duplicates inverted repeats of variable length and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=PG18;
Calcutt M.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=9911553; PubMed=9916088; Calcutt M.J., Kim M.F., Karpas A.B., Muhlradt P.F., Wise K.S.; Calcutt M.J., Kim M.F., Karpas A.B., Muhlradt p.F., Wise K.S.; "Differential posttranslational processing confers intraspecies variation of a major surface lipoprotein and a macrophage-activating lipopeptide of Mycoplasma fermentans."; Infect. Immun. 67:760-771(1999).
                                                                                                                                        METAL
                                                                                                                                                      NON_TER
                                                                                                                                                                                                    PFAM; PF00131; metalthio;
                                                                                                                                                                                                                                                                                                                                           Huang
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sus scrota (Pig).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                               Metal-binding;
                                                                                                                                                                             PROSITE;
                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                         INTERPRO;
                                                                                                                                                                                                                                                                                                                              duang M.C., Pan P.K., Zeng T.-F., Cher
Multiple isoforms of metallothionein
                                                                                                                                                                                                                                                                                                                                                                    TISSUE=LIVER;
                                                                                                                                                                                                              NTERPRO;
                                                                                                                                                                                                                                                                                                                     iver
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nes 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CXXXCXXXXXXXXXXXXXX 20
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                                                                                                                                                                                                                                                            GLUCOCORTICOIDS
                                                                                                                                                                                                                                                                    FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE RESIDUES THAT BIND VARIOUS HEAVY METALS; THESE PROTEINS ARE TRANSCRIPTIONALLY REGULATED BY BOTH HEAVY METALS AND
                                                                                                                                                                                                                                                                                            > 211:49-55(1998)
FUNCTION: METALL
                                                                                                                                                                                      PR00860; MTVERTEBRATE
                                                                                                                                                                         PS00203;
                                                                                                                                                                                                              IPR000006; -. IPR003019; -.
                                                                   48
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         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                               Metal-thiolate cluster;
                                                                                                                                                                                                                                                BAA19179.1;
                                                                                                                                                                          METALLOTHIONEIN_VRT;
                                                                                                                                                                                                                                                                                                                                                    PubMed=9573337;
                                                                   4704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5319 MW;
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                   100.0%
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                                                                   MW;
                              08;
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         16;
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                   Pred. No.
                                                                                         CLUSTER.
CLUSTER.
CLUSTER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Created)
                                                                   CLUSTER.
F0F5AC96BB9EBD19 CRC64;
                                                                                                                           CLUSTER
                                                                                                                                        CLUSTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        668836FA3592B2C7 CRC64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 52;
         Mismatches
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                              52;
                                                                                                                                                                                                                                                                                                                               are
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                                                                                                                                                               1.
Chelation; Zinc
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                               DB 6;
                                                                                                                                                                                                                                                                                                                              expressed in
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         0;
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                              Length 48;
        Indels
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Sus.
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Best Local S
Matches 4
                                                                                                                                                                                                        Q23947;
01-NOV-1996
01-NOV-1996
01-OCT-2000
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01-MAY-1997
01-OCT-2000
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SEQUENCE
                                                                                                                                                                                     01-NOV-1996 (TrEMBLrel.
01-NOV-1996 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
TESTIS-SPECIFIC RNA.
                                                                                                                                                                                                                                                                                             Q23947
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Huang M.C., Pan P.K., Zeng T.-F., Chen N.C., Peng J.Y., Huang P.C.,
"Multiple isoforms of metallothionein are expressed in the porcine
                       SEQUENCE FROM N.A.
                                                              NCBI_TaxID=7224;
                                                                                                       Pterygota;
                                                                                                                             Eukaryota;
                                                                                                                                                Drosophila
                                                                                                                                                                      DHTC3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PFAM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INTERPRO; IPR003019;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
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HSSP; P02795; 2MHU.
INTERPRO; IPR000006; -.
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E 211:49-55(1998).

E 211:49-55(1998).

FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE FUNCTIONS THAT BIND VARIOUS HEAVY METALS; THESE PROTEINS ARE RESIDUES THAT BIND VARIOUS HEAVY METALS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G
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                                                                               Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Neoptera; Endopterygota; Diptera; Brachycera; Musc
a; Drosophilidae; Drosophila.
                                                                                                                                                hydei (Fruit fly)
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(TrEMBLrel. 03, Last sequence update)
(TrEMBLrel. 15, Last annotation update)
                                                                                                                                                                                                                                                                                               PRELIMINARY;
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Last annotation updat
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Pred. No. 80;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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Best Local Similarity 20.0
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
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Q23948;
01-NOV-1996 (TrEMBLTel. (
01-NOV-1996 (TrEMBLTel. (
01-OCT-2000 (TrEMBLTel. 1
TESTIS-SPECIFIC RNA.
SEQUENCE FROM GLAND;
TISSUE=VENOM GLAND;
TISSUE=VENOM GLAND;
MEDLINE=99058072; PubMed=9837992;
Liu L.F., Chang C.C., Liau M.Y., Kuo K.W.;
Liu L.F., Chang C.C., Liau M.Y., Kuo K.W.;
"Genetic characterization of the mRNAs encoding alpha-bungarotoxin: isoforms and RNA editing in Bungarus multicinctus gland cells.";
isoforms and RNA editing in Bungarus multicinctus gland cells.";
                                                                                                                                                                                                                                                                                                                                                                                                                  0]YLIU;
01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
ALPHA-BUNGAROTOXIN ISOFORM R7 (FRAGMENT).
Bungarus multicinctus (Many-banded krait).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrat
Eukaryota; Metazoa; Squamata; Scieroglossa; Serpentes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q9YI10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kuech A., Buenemann H.;
Submitted (DEC-1990) to the EMBL/GenBank/DDBJ
EMBL; X58111; CAA41118.1; -.
FLYBASE; FB9n0015097; Dhyd\Mst87Fb.
INTERPRO; IPR001450; -.
PROSITE; PS00198; 4FE4S; FERREDOXIN; UNKNOWN_1.
SEQUENCE 49 AA; 4550 MW; 24758FEC51369FID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; MuscEphydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (DEC-1990) to the EMBL/Gen EMBL; X58114: CAAA1117.1; -. FLYBASE; FBGn0015096; Dhyd\mst87Fa. INTERPRO; IPR001450; -. PROSITE; PS00198; 4FE4S_FERREDOXIN; SEQUENCE 49 AA; 4610 MW; BE158FE
                                                                                                                                                                                                                                                                                                                                                                 Elapidae; Bungarinae;
NCBI_TaxID=8616;
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4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                          Bungarus.
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P82107;
                      Eur.
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                                                                                               Moser M., Auerswald E., "Bdellastatin, a serine from the medical leech
                                                                                                                                                                                                                Hirudo medicinalis (Medicinal leech).
Eukaryota; Metazoa; Annelida; Clitellata; Hirudinida;
Arynchobdellida; Hirudiniformes; Hirudinidae; Hirudo.
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INTERPRO; IPR000027; -.
                                                                                                                                                         SEQUENCE.
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"Characterisation of metallothioneins sequenies in three clams
Ruditapes decussatus, Ruditapes philippinarum and Venerupis
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Veneroidea; Veneridae; Ruditap
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57
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STRONG INHIBITOR OF MAMMALIAN
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                                                                        E., Mentele R., Eckerskorn C., Fritz H., Fink E ine protease inhibitor of the antistasin family ch (Hirudo medicinalis). Primary structure, and characterization of native and recombinant
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01-OCT-2000 (TrEMBLrel. 15, L
01-OCT-2000 (TrEMBLrel. 15, L
METALLOTHIONEIN (FRAGMENT).
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      Cyprinodon sp.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Ecukaryota; Metazoa; Chordata; Teleostei; Euteleostei; Neoteleostei;
Actinopterygii; Neopterygii; Percomorpha; Atherinomorpha;
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DISULFID
SEQUENCE
Cyprinodontiformes; Cyprinodontidae;
                                              MT A.
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Moraga D. Tanguy A.;
"Characterisation of metallothioneins sequenies
"Characterisation, Ruditapes philippinarum ar
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Eukaryota; Metazoa; Mollusca; Bivalvia;
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01-AUG-1998
01-OCT-2000
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PROSITE; PS00203; METALLOTHIONEIN VRT;
PROSITE; PS00203; METALLOTHIOLATE cluster;
EMBL;
                        "Metallothionein in Antarctic organisms.";
Submitted (SEP-1998) to the EMBL/GenBank/DBJ databases.
-!- PUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE RESIDUES THAT BIND VARIOUS HEAVY METALS; THESE PROTEINS ARE TRANSCRIPTIONALLY REGULATED BY BOTH HEAVY METALS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                 METAL
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HSSP;
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SEQUENCE FROM N.A.
                                                                                                                         Scudiero R., Verde C., Parisi E.;
                                                                                                                                                            Notothenioidei; NCBI_TaxID=8209,
                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata;
Actinopterygii; Neopterygii; Teleostei;
                                                                                                                                                                                                          and Chionodraco rastrospinosus.
                                                                                                                                                                                                                                       METALLOTHIONEIN
                                                                                                                                                                                                                                                                                      073914
                                                                         Carginale V., Capasso
                                                                                    TISSUE-LIVER;
                                                                                                               Submitted (JUN-1998) to
                                                                                                                        Parisi
                                                                                                                                           SEQUENCE FROM N.A
                                                                                                                                                                                Acanthomorpha;
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nes 4; Conser
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mitted (APR-1996) to the EMBL/GenBank/DDBJ databases.
FUNCTION: METALLOTHIOMEINS HAVE A HIGH CONTENT OF CYSTEINE
RESIDUES THAT BIND VARIOUS HEAVY METALS; THESE PROTEINS ARE
                   GLUCOCORTICOIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GLUCOCORTICOIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TRANSCRIPTIONALLY REGULATED BY BOTH HEAVY METALS AND
                                                                                                                                                                                                                                                                                                                                    CKKSCCSCCPSGCSKCASGC
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AJ006484;
AJ011584;
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CAA07063.
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Euteleostei; Neoteleostei;
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PROSTTE; PS00203; METALLOTHIONEIN_VRT; 1.
Metal-binding; Metal-thiolate cluster; Ch
METAL 23 CLUSTER.
METAL 26 26 CLUSTER.
METAL 16 16 CLUSTER.
METAL 18 18 CLUSTER.
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METAL 21 21 CLUSTER.
METAL 30 30 CLUSTER.
                                                                                                                                                                                                                                       Carginale V., Capasso C., Scudiero R., Parisi E.;
"Metallothionein in Antarctic organisms.";
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE RESIDUES THAT BIND VARIOUS HEAVY METALS; THESE PROTEINS ARI
                                                                                     METAL
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                                                                                                                                                                                                                                                                                                                                      Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Notothenioidei; Channichthyidae; Chaenocephalus.
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Eukaryota; Metazoa; Chordata;
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01-NOV-1998 (TrEMBLrel.
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01-OCT-2000 (TrEMBLrel.
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PROSITE; PS00203; METALLOTHIONEIN_VRT;
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No. 91;
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Teleostei;
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Best Local S
Matches 4
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PROSITE; PS00203; METALLOTHIONEIN_VRT; 1.
Metal-binding; Metal-thiolate cluster; Ch
METAL 26 26 CLUSTER.
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METAL 16 16 CLUSTER.
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TISSUE-LIVER;

Carginale V., Capasso C., Scudiero R., Parisi E.;

"Metallothionein in Antarctic organisms.";

"Metallothionein in Antarctic organisms.";

Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases.

-i- FUNCTION: METALLOTHIONEINS HAVE A HIGH CONTENT OF CYSTEINE
RESIDUES THAT BIND VARIOUS HEAVY METALS; THESE PROTEINS ARE
TERMIQUES THAT BIND VARIOUS HEAVY METALS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         O93609 PRELIMINARY; PRT; 60 AA.
093609;
01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei

Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes;

Notothenioidei; Nototheniidae; Trematomus.
                                                                                                                                                                                                                                                                                                                                          EMBL; AJ011585; CAA09715.1; HSSP; P02802; 1DFS. INTERPRO; IPR000006; -. INTERPRO; IPR003019; -. PFAM; PF00131; metalthio; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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                                                                   l Similarity
4; Conserv
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16; Mismatches
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AC66F3015A2C4251 CRC64;
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